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RAW SEQUENCE LISTING  
PATENT APPLICATION: US/09/723,676

DATE: 12/08/2000  
TIME: 15:22:00

Input Set : A:\001214seq.txt  
Output Set: N:\CRF3\12082000\I723676.raw

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3 <110> APPLICANT: Welcher, Andrew A.
4           Jing, Shuqian
5           Luethy, Roland
7 <120> TITLE OF INVENTION: Interleukin-1 Receptor Antagonist-Like Molecules and
8           Uses Thereof
10 <130> FILE REFERENCE: 00-1214
C--> 12 <140> CURRENT APPLICATION NUMBER: US/09/723,676
C--> 13 <141> CURRENT FILING DATE: 2000-11-28
15 <150> PRIOR APPLICATION NUMBER: 60/170,052
16 <151> PRIOR FILING DATE: 1999-12-10
18 <160> NUMBER OF SEQ ID NOS: 18
20 <170> SOFTWARE: PatentIn Ver. 2.0
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24 <212> TYPE: DNA
25 <213> ORGANISM: Homo sapiens
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29 <222> LOCATION: (301)..(774)
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36 cctggagtc ctggaccgc tttcatgca gtagaccac ttacatgcaa cttaggcct 180
38 gactttgaaa cgaggcacaa aagcatctct tgctaaagggt aacttctgct gcttagaacc 240
40 cagcctctc accaccatct gatctatctt gttctcttca caaaaggqctc tqaayacatc 300
42 atg aac cca caa cgg gag gca ccc aaa tcc tat gct att cgt gat 348
43 Met Asn Pro Gln Arg Glu Ala Ala Pro Lys Ser Tyr Ala Ile Arg Asp
44      1          5          10          15
46 tct cga cag atg gtg tgg gtc ctg agt gga aat tct tta ata gca gct 396
47 Ser Arg Gln Met Val Trp Val Leu Ser Gly Asn Ser Leu Ile Ala Ala
48      20          25          30
50 cct ctt agc cgc agc att aag oct gtc act ctt cat tta ata gcc tgt 444
51 Pro Leu Ser Arg Ser Ile Lys Pro Val Thr Leu His Leu Ile Ala Cys
52      35          40          45
54 aga gac aca gaa ttc agt gac aag gaa aag ggt aat atg gtt tac ctg 492
55 Arg Asp Thr Glu Phe Ser Asp Lys Glu Lys Gly Asn Met Val Tyr Leu
56      50          55          60
58 gga atc aag gga aaa gat ctc tgt ctc ttc tgt gca gaa att cag ggc 540
59 Gly Ile Lys Gly Lys Asp Leu Cys Leu Phe Cys Ala Glu Ile Gln Gly
60      65          70          75          80
62 aag cct act ttg cag ctt aag gaa aaa aat atc atg gac ctg tat gtg 588
63 Lys Pro Thr Leu Gln Leu Lys Glu Lys Asn Ile Met Asp Leu Tyr Val
64      85          90          95
66 gag aag aaa gca cag aag ccc ttt ctc ltt ttc cac aat aaa gaa gyc 636
67 Glu Lys Lys Ala Gln Lys Pro Phe Leu Phe Phe His Asn Lys Glu Gly
68      100         105         110
70 tcc act tct gtc ltt cag tca gtc tct tac cct ggc tgg ttc ata gcc 684

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71 Ser Thr Ser Val Phe Gln Ser Val Ser Tyr Pro Gly Trp Phe Ile Ala
72      115          120          125
74 acc tcc acc aca tca gga caq ccc atc itt ctc acc aag gaa aga ggc  732
75 Thr Ser Thr Ser Gly Gln Pro Ile Phe Leu Thr Lys Glu Arg Gly
76      130          135          140
78 ata act aat aac act aac ttc tac tta gat tct gtg gaa taa  774
79 Ile Thr Asn Asn Ile Asn Phe Tyr Leu Asp Ser Val Glu
80 145          150          155
82 atccagccta ggctgtgggt ggctgggtcc agqatacgaga atcaagctgt cagagtcata 834
84 ttaacagatc attatgcgc ttagttcaact agcaatgcac cccatccata gtttacccca 894
86 ttcttactat ccaaaggcca ctctctccca caaacatcca ttctgtacc aagaccctca 954
88 ctcaatgtc actatccaa qatqaaaacct aaaaatcaact ttccattttt tcttgatott 1014
90 acccccacccat ccactcaatgc gccatgcccc gtttagttaa ccccccaaat gctgcttcat 1074
92 gcaacccatcc attccatttc ctttgtccaa cccatgtatq agagatgtgg attcatqaca 1134
94 ttttggatcat acaacttccat caataaaaca ttataatatg tgccccaaag ataaagctga 1194
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101 <212> TYPE: PRT
102 <213> ORGANISM: Homo sapiens
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109      20          25          30
111 Pro Leu Ser Arg Ser Ile Lys Pro Val Thr Leu His Leu Ile Ala Cys
112      35          40          45
114 Arg Asp Thr Glu Phe Ser Asp Lys Glu Lys Gly Asn Met Val Tyr Leu
115      50          55          60
117 Gly Ile Lys Gly Lys Asp Leu Cys Leu Phe Cys Ala Glu Ile Gln Gly
118      65          70          75          80
120 Lys Pro Thr Leu Gln Leu Lys Glu Lys Asn Ile Met Asp Leu Tyr Val
121      85          90          95
123 Glu Lys Lys Ala Gln Lys Pro Phe Leu Phe His Asn Lys Glu Gly
124      100         105         110
126 Ser Thr Ser Val Phe Gln Ser Val Ser Tyr Pro Gly Trp Phe Ile Ala
127      115         120         125
129 Thr Ser Thr Ser Gly Gln Pro Ile Phe Leu Thr Lys Glu Arg Gly
130      130         135         140
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133 145          150          155
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138 <212> TYPE: PRT
139 <213> ORGANISM: Homo sapiens
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142 Met Asn Pro Gln Arg Glu Ala Ala Pro Lys Ser Tyr Ala Ile Arg Asp
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146               20               25               30  
 148 Pro Leu Ser Arg Ser Ile Lys Pro Val Thr Leu His Leu Ile Ala Cys  
 149               35               40               45  
 151 Arg Asp Thr Glu Phe Ser Asp Lys Glu Lys Gly Asn Met Val Tyr Leu  
 152               50               55               60  
 154 Gly Ile Lys Gly Lys Asp Leu Cys Leu Phe Cys Ala Glu Ile Gln Gly  
 155               65               70               75               80  
 157 Lys Pro Thr Leu Gln Leu Lys Leu Gln Gly Ser Gln Asp Asn Ile Gly  
 158               85               90               95  
 160 Lys Asp Thr Cys Trp Lys Leu Val Gly Ile His Thr Cys Ile Asn Leu  
 161               100              105              110  
 163 Asp Val Arg Glu Ser Cys Phe Met Gly Thr Leu Asp Gln Trp Gly Ile  
 164               115              120              125  
 166 Gly Val Gly Arg Lys Lys Trp Lys Ser Ser Phe Gln His His His Leu  
 167               130              135              140  
 169 Arg Lys Lys Asp Lys Asp Phe Ser Ser Met Arg Thr Asn Ile Gly Met  
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 178 <212> TYPE: PRT  
 179 <213> ORGANISM: Homo sapiens  
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 183     1              5              10              15  
 185 Asp Ile Asn His Arg Val Trp Val Leu Gln Asp Gln Thr Leu Ile Ala  
 186     20              25              30  
 188 Val Pro Arg Lys Asp Arg Met Ser Pro Val Thr Ile Ala Leu Ile Ser  
 189     35              40              45  
 191 Cys Arg His Val Glu Thr Leu Glu Lys Asp Arg Gly Asn Pro Ile Tyr  
 192     50              55              60  
 194 Leu Gly Leu Asn Gly Leu Asn Leu Cys Leu Met Cys Ala Lys Val Gly  
 195     65              70              75              80  
 197 Asp Gln Pro Thr Leu Gln Leu Lys Glu Lys Asp Ile Met Asp Leu Tyr  
 198     85              90              95  
 200 Asn Gln Pro Glu Pro Val Lys Ser Phe Leu Phe Tyr His Ser Gln Ser  
 201     100              105              110  
 203 Gly Arg Asn Ser Thr Phe Glu Ser Val Ala Phe Pro Gly Trp Phe Ile  
 204     115              120              125  
 206 Ala Val Ser Ser Glu Gly Gly Cys Pro Leu Ile Leu Thr Gln Glu Leu  
 207     130              135              140  
 209 Gly Lys Ala Asn Thr Thr Asp Phe Gly Leu Thr Met Leu Phe  
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 215 <212> TYPE: PRT  
 216 <213> ORGANISM: Homo sapiens  
 218 <400> SEQUENCE: 5  
 219 Met Glu Ile Cys Arg Gly Leu Arg Ser His Leu Ile Thr Leu Leu

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220   1          5          10          15
222 Phe Leu Phe His Ser Glu Thr Ile Cys Arg Pro Ser Gly Arg Lys Ser
223           20          25          30
225 Ser Lys Met Gln Ala Phe Arg Ile Trp Asp Val ASN Gln Lys Thr Phe
226           35          40          45
228 Tyr Leu Arg Asn Asn Gln Leu Val Ala Gly Tyr Leu Gln Gly Pro Asn
229           50          55          60
231 Val Asn Leu Glu Glu Lys Ile Asp Val Val Pro Ile Glu Pro His Ala
232           65          70          75          80
234 Leu Phe Leu Gly Ile His Gly Gly Lys Met Cys Leu Ser Cys Val Lys
235           85          90          95
237 Ser Gly Asp Glu Thr Arg Leu Gln Leu Glu Ala Val Asn Ile Thr Asp
238           100         105         110
240 Leu Ser Glu Asn Arg Lys Gln Asp Lys Arg Phe Ala Phe Ile Arg Ser
241           115         120         125
243 Asp Ser Gly Pro Thr Thr Ser Phe Glu Ser Ala Ala Cys Pro Gly Trp
244           130         135         140
246 Phe Leu Cys Thr Ala Met Glu Ala Asp Gln Pro Val Ser Leu Thr Asn
247           145         150         155         160
249 Met Pro Asp Glu Gly Val Met Val Thr Lys Phe Tyr Phe Gln Glu Asp
250           165         170         175
252 Glu
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258 <212> TYPE: PRT
259 <213> ORGANISM: Homo sapiens
261 <400> SEQUENCE: 6
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263   1          5          10          15
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266           20          25          30
268 Gly Gln Asp Met Glu Gln Gln Val Val Phe Ser Met Ser Phe Val Gln
269           35          40          45
271 Gly Glu Glu Ser Asn Asp Lys Ile Pro Val Ala Leu Gly Leu Lys Glu
272           50          55          60
274 Lys Asn Leu Tyr Leu Ser Cys Val Leu Lys Asp Asp Lys Pro Thr Leu
275           65          70          75          80
277 Gln Leu Glu Ser Val Asp Pro Lys Asn Tyr Pro Lys Lys Lys Met Glu
278           85          90          95
280 Lys Arg Phe Val Phe Asn Lys Ile Gln Ile Asn Asn Lys Leu Glu Phe
281           100         105         110
283 Glu Ser Ala Gln Phe Pro Asn Trp Tyr Ile Ser Thr Ser Gln Ala Glu
284           115         120         125
286 Asn Met Pro Val Phe Leu Gly Gly Thr Lys Gly Gln Asp Ile Thr
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289 Asp Phe Thr Met Gln Phe Val Ser Ser
290           145         150
293 <210> SEQ ID NO: 7
294 <211> LENGTH: 11

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295 <212> TYPE: PRT  
296 <213> ORGANISM: Human immunodeficiency virus type 1  
298 <400> SEQUENCE: 7  
299 Tyr Gly Arg Lys Lys Arg Arg Gln Arg Arg Arg  
300 1 5 10  
303 <210> SEQ ID NO: 8  
304 <211> LENGTH: 15  
305 <212> TYPE: PRT  
306 <213> ORGANISM: Artificial Sequence  
308 <220> FEATURE:  
309 <223> OTHER INFORMATION: Description of Artificial Sequence: internalizing  
310 domain derived from HIV tat protein  
312 <400> SEQUENCE: 8  
313 Gly Gly Gly Gly Tyr Gly Arg Lys Lys Arg Arg Gln Arg Arg Arg  
314 1 5 10 15  
317 <210> SEQ ID NO: 9  
318 <211> LENGTH: 23  
319 <212> TYPE: DNA  
320 <213> ORGANISM: Artificial Sequence  
322 <220> FEATURE:  
323 <223> OTHER INFORMATION: Description of Artificial Sequence:  
324 Oligonucleotide 2362-94  
326 <400> SEQUENCE: 9  
327 catggacctg tatgtggaga aga 23  
330 <210> SEQ ID NO: 10  
331 <211> LENGTH: 23  
332 <212> TYPE: DNA  
333 <213> ORGANISM: Artificial Sequence  
335 <220> FEATURE:  
336 <223> OTHER INFORMATION: Description of Artificial Sequence:  
337 Oligonucleotide 2362-95  
339 <400> SEQUENCE: 10  
340 gccagggttaa gagactgact gaa 23  
343 <210> SEQ ID NO: 11  
344 <211> LENGTH: 23  
345 <212> TYPE: DNA  
346 <213> ORGANISM: Artificial Sequence  
348 <220> FEATURE:  
349 <223> OTHER INFORMATION: Description of Artificial Sequence:  
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352 <400> SEQUENCE: 11  
353 aqcgguataac aatttcacac agg 23  
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358 <212> TYPE: DNA  
359 <213> ORGANISM: Artificial Sequence  
361 <220> FEATURE:  
362 <223> OTHER INFORMATION: Description of Artificial Sequence:  
363 Oligonucleotide 2366-21

VERIFICATION SUMMARY  
PATENT APPLICATION: US/09/723,676 DATE: 12/08/2000  
TIME: 15:22:01

Input Set : A:\001214seq.txt  
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L:13 M:271 C: Current Filing Date differs, Replaced Current Filing Date